

### **IN THE SPECIFICATION**

On page 6, line 4 replace the 1<sup>st</sup> paragraph with the following:

- -Figure 5 shows an amino acid sequence alignment for the PHD finger motif of AIR-1 (SEQ ID NO:38, Mi-2 SEQ ID NO:39 and SEQ ID NO:40), AND TIF1 (SEQ ID NO:41). The consensus amino acid residues conserved in the PHD finger motif is indicated by the bold letters underneath. The residues that are identical with AIR-1 (aa 299-340) (SEQ. ID NO: 37) are shown by the dots. GenBank accession nos. of Mi-2 and TIF1 are X86691 and AF009353, respectively.- -